

SEQUENCE LISTING

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      Acoustic Absorption Polymers and Their Methods of Use
      BERL025/01US
<130>
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 Gly Gly Phe Pro
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Gly Gly Ala Pro
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      Residue at position 4 is V, E, F or I
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Gly Xaa Gly Xaa Pro
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<210> 7
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Ala Pro Gly Val Gly Val
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      (1)..(35)
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Gly Val Gly Val Pro Gly Val Gly Phe Pro Gly Glu Gly Phe Pro Gly
Val Gly Val Pro Gly Val Gly Phe Pro Gly Phe Gly Phe Pro Gly Val
                                25
Gly Val Pro
        35
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       (1)..(35)
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<400> 9
Gly Val Gly Val Pro Gly Val Gly Phe Pro Gly Glu Gly Phe Pro Gly
Val Gly Val Pro Gly Val Gly Phe Pro Gly Val Gly Phe Pro Gly Val
                                25
Gly Val Pro
        35
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<221> PEPTIDE
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<400> 10
Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Glu Gly Val Pro Gly
                5
                                                         15
Val Gly Val Pro Gly Val Gly Phe Pro Gly Phe Gly Phe Pro Gly Val
                                 25
Gly Val Pro
        35
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       (1)..(35)
<400> 11
Gly Val Gly Val Pro Gly Val Gly Phe Pro Gly Glu Gly Phe Pro Gly
                5
1
                                                         15
Val Gly Val Pro Gly Val Gly Val Pro Gly Val Pro Gly Val
            20
Gly Val Pro
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<400> 12
Gly Val Gly Val Pro Gly Val Pro Gly Glu Gly Val Pro Gly
                                     10
Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val
                                25
Gly Val Pro
        35
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       (1)..(65)
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Gly Val Gly Ile Pro Gly Phe Gly Glu Pro Gly Glu Gly Phe Pro Gly
Val Gly Val Pro Gly Phe Gly Phe Pro Gly Phe Gly Ile Pro Gly Val
            20
                                 25
Gly Ile Pro Gly Phe Gly Glu Pro Gly Glu Gly Phe Pro Gly Val Gly
Val Pro Gly Phe Gly Phe Pro Gly Phe Gly Ile Pro Gly Val Gly Val
Pro
65 ·
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Gly Val Gly Val Pro Gly Val Gly Phe Pro Gly Lys Gly Phe Pro Gly
Val Gly Val Pro Gly Val Gly Phe Pro Gly Phe Gly Phe Pro Gly Val
            20
                                 25
                                                      30
Gly Val Pro
        35
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Gly Val Gly Val Pro Gly Val Gly Phe Pro Gly Lys Gly Phe Pro Gly
Val Gly Val Pro Gly Val Gly Phe Pro Gly Val Gly Phe Pro Gly Val
            20
                                                      30
Gly Val Pro
        35
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PEPTIDE
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<400>
       16
Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Lys Gly Val Pro Gly
                5
                                     10
                                                          15
Val Gly Val Pro Gly Val Gly Phe Pro Gly Phe Gly Phe Pro Gly Val
                                 25
Gly Val Pro
        35
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       (1)..(35)
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       17
Gly Val Gly Val Pro Gly Val Gly Phe Pro Gly Lys Gly Phe Pro Gly
                5
1
                                     10
                                                          15
Val Gly Val Pro Gly Val Gly Val Pro Gly Val Pro Gly Val
            20
                                                      30
Gly Val Pro
        35
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       18
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       (1)..(35)
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<400> 18
Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Lys Gly Val Pro Gly
Val Gly Val Pro Gly Val Gly Val Pro Gly Val Pro Gly Val
                                 25
Gly Val Pro
        35
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       35
       PRT
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       (1)..(35)
<400> 19
Gly Val Gly Val Pro Gly Val Gly Phe Pro Gly Glu Gly Phe Pro Gly
Val Gly Val Pro Gly Val Gly Phe Pro Gly Lys Gly Val Pro Gly Val
                                 25
                                                     30
            20
Gly Val Pro
        35
<210>
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       (1)..(35)
<400>
       20
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Gly Val Gly Val Pro Gly Val Gly Phe Pro Gly Glu Gly Phe Pro Gly
Val Gly Val Pro Gly Val Gly Val Pro Gly Lys Gly Val Pro Gly Val
                                25
Gly Val Pro
       35
<210>
      21
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      5
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<222> (4)..(4)
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      Residue at position 4 is modified to have
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<400> 21
Val Pro Gly Xaa Gly
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Ile Pro Gly Val Gly
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<211> 11
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      Residue at position 6 is S, T or Y
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Gly Val Gly Val Pro Xaa Gly Val Gly Val Pro
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      Residue at position 2 is V, E, F, Y, K, S or T
      Residue at position 4 is V, E, F, I, S, T or Y
      Al least one of residues at positions 2 or 4 is S, T or Y
<400> 24
Gly Xaa Gly Xaa Pro
<210>
      25
<211>
      30
<212> PRT
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<220>
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       (1)..(30)
<400> 25
Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Lys Gly Val Pro Gly
                5
                                    10
                                                         15
Val Gly Val Pro Gly Val Gly Phe Pro Gly Phe Gly Phe Pro
            20
                                25
                                                     30
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      (1)..(66)
       26
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gaggatccag gcgttggggt accgggtgtt ggcgatccgg gtaaaggtgt cccggggttg
gtgtgc
   66
<210>
       27
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       66
<212> DNA
<213> Artificial
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<221> misc_structure
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      (1)..(66)
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ctggatccaa cgcctgggaa tccgaaaccc ggaaagccta cacccggcac accaacgccc
   60
gggaca
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66

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<210>
       28
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       (1)..(10)
<400>
       28
Gly Val Gly Val Pro Gly Tyr Gly Val Pro
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       PEPTIDE
<222>
      (1)..(45)
<400>
       29
Gly Val Gly Ile Pro Gly Glu Gly Ile Pro Gly Val Gly Ile Pro Gly
                                                          15
                                     10
Val Gly Ile Pro Gly Glu Gly Ile Pro Gly Val Gly Ile Pro Gly Val
Gly Ile Pro Gly Glu Gly Ile Pro Gly Val Gly Ile Pro
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<211>
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Gly Val Gly Ile Pro Gly Val Gly Ile Pro Gly Glu Gly Ile Pro Gly
Val Gly Ile Pro Gly Val Gly Ile Pro Gly Val Gly Ile Pro
<210> 31
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Gly Glu Gly Ile Pro Gly Val Gly Ile Pro Gly Glu Gly Ile Pro Gly
                                    10
Val Gly Ile Pro Gly Val Gly Ile Pro Gly Val Gly Ile Pro
            20
                                25
                                                    30
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       32
<211> 45
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<222> (1)..(45)
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Gly Val Gly Ile Pro Gly Lys Gly Ile Pro Gly Val Gly Ile Pro Gly
Val Gly Ile Pro Gly Lys Gly Ile Pro Gly Val Gly Ile Pro Gly Val
                                 25
Gly Ile Pro Gly Lys Gly Ile Pro Gly Val Gly Ile Pro
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       30
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       (1)..(30)
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       33
Gly Val Gly Ile Pro Gly Val Gly Ile Pro Gly Lys Gly Ile Pro Gly
Val Gly Ile Pro Gly Val Gly Ile Pro Gly Val Gly Ile Pro
            20
                                 25
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       30
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       (1)..(30)
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       34
Gly Lys Gly Ile Pro Gly Val Gly Ile Pro Gly Lys Gly Ile Pro Gly
                5
1
                                     10
                                                          15
Val Gly Ile Pro Gly Val Gly Ile Pro Gly Val Gly Ile Pro
```

30

20

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25
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       35
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       (1)..(110)
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Gly Val Gly Ile Pro Gly Val Gly Ile Pro Gly Val Gly Ile Pro Gly
Val Gly Ile Pro Gly Val Gly Ile Pro Gly Val Gly Ile Pro Gly Val
                                 25
Gly Ile Pro Gly Val Gly Ile Pro Gly Val Gly Ile Pro Gly Val Gly
Ile Pro Gly Val Gly Ile Pro Gly Val Gly Ile Pro Gly Val Gly Ile
    50
                         55
                                              60
Pro Gly Val Gly Ile Pro Gly Val Gly Ile Pro Gly Val Gly Ile Pro
65
                     70
                                          75
                                                              80
Gly Val Gly Ile Pro Gly Val Gly Ile Pro Gly Val Gly Ile Pro Gly
                85
Val Gly Ile Pro Gly Val Gly Ile Pro Gly Tyr Gly Ile Pro
            100
                                 105
                                                      110
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Residue at position 107 is associated with an SO4 ion
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       36
Gly Val Gly Ile Pro Gly Val Gly Ile Pro Gly Val Gly Ile Pro Gly
                                                         15
Val Gly Ile Pro Gly Val Gly Ile Pro Gly Val Gly Ile Pro Gly Val
Gly Ile Pro Gly Val Gly Ile Pro Gly Val Gly Ile Pro Gly Val Gly
Ile Pro Gly Val Gly Ile Pro Gly Val Gly Ile Pro Gly Val Gly Ile
Pro Gly Val Gly Ile Pro Gly Val Gly Ile Pro Gly Val Gly Ile Pro
Gly Val Gly Ile Pro Gly Val Gly Ile Pro Gly Val Gly Ile Pro Gly
                                                         95
Val Gly Ile Pro Gly Val Gly Ile Pro Gly Tyr Gly Ile Pro
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            100
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Gly Val Gly Ile Pro Gly Val Gly Ile Pro Gly Val Gly Ile Pro Gly
Val Gly Ile Pro Gly Val Gly Ile Pro Gly Val Gly Ile Pro Gly Val
            20
                                25
Gly Ile Pro Gly Val Gly Ile Pro Gly Val Gly Ile Pro Gly Val Gly
        35
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Ile Pro Gly Val Gly Ile Pro Gly Tyr Gly Ile Pro
                         55
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       38
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       60
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       Residue at position 58 is associated with an SO4 ion
<400>
       38
Gly Val Gly Ile Pro Gly Val Gly Ile Pro Gly Val Gly Ile Pro Gly
                5
                                      10
                                                          15
Val Gly Ile Pro Gly Val Gly Ile Pro Gly Val Gly Ile Pro Gly Val
            20
Gly Ile Pro Gly Val Gly Ile Pro Gly Val Gly Ile Pro Gly Val Gly
                             40
        35
Ile Pro Gly Val Gly Ile Pro Gly Tyr Gly Ile Pro
                                              60
    50
                         55
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       39
Gly Val Gly Ile Pro Gly Val Gly Ile Pro Gly Val Gly Ile Pro Gly
                5
                                      10
                                                          15
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Val Gly Ile Pro Gly Val Gly Ile Pro Gly Val Gly Ile Pro Gly Val
Gly Ile Pro Gly Val Gly Ile Pro Gly Tyr Gly Ile Pro
                             40
<210>
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<400> 40
Gly Val Gly Ile Pro Gly Val Gly Ile Pro Gly Val Gly Ile Pro Gly
                                     10
Val Gly Ile Pro Gly Val Gly Ile Pro Gly Val Gly Ile Pro Gly Val
                                 25
Gly Ile Pro Gly Val Gly Ile Pro Gly Tyr Gly Ile Pro
                             40
        35
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       41
Gly Val Gly Ile Pro Gly Val Gly Ile Pro Gly Val Gly Ile Pro Gly
1
                5
                                     10
                                                          15
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Val Gly Ile Pro Gly Val Gly Ile Pro Gly Tyr Gly Ile Pro

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This is a synthetic sequence.

<223>

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      44
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       45
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       10
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      (1)..(10)
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Ile Pro Gly Val Gly Ile Pro Gly Tyr Gly
                5
                                     10
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       46
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       10
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Ile Pro Gly Val Gly Ile Pro Gly Tyr Gly
                                     10
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<210> 47
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<223> This is a synthetic sequence.
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Ile Pro Gly Val Gly Ile Pro Gly Tyr Gly
1 5 10
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